

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/550,797
Source: IFW
Date Processed by STIC: 8/28/06

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 08/28/2006

PATENT APPLICATION: US/10/550,797

TIME: 09:26:32

Input Set : F:\seq_210121_609uspc.app.txt

Output Set: N:\CRF4\08282006\J550797.raw

3 <110> APPLICANT: Zehentner-Wilkinson, Barbara K.
 4 Hayes, Dawn
 5 Houghton, Raymond L.
 7 <120> TITLE OF INVENTION: METHODS, COMPOSITIONS AND KITS FOR THE DETECTION
 8 AND MONITORING OF LUNG CANCER
 10 <130> FILE REFERENCE: 210121.609USPC
 12 <140> CURRENT APPLICATION NUMBER: US 10/550,797
 C--> 13 <141> CURRENT FILING DATE: 2005-09-22
 15 <160> NUMBER OF SEQ ID NOS: 34
 17 <170> SOFTWARE: Corixa Invention Disclosure Database
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 3951
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Homo sapiens
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 37 ctgtattatt agtaagcttt ttaaagaagg atgcacctt atctacaata gcacccaaaa 780
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92 <210> SEQ ID NO: 2

93 <211> LENGTH: 943

94 <212> TYPE: PRT

95 <213> ORGANISM: Homo sapiens

97 <400> SEQUENCE: 2

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100 Thr Leu Leu Val Ala Leu Ser Ser Glu Leu Pro Phe Leu Gly Ala Gly
101 20 25 30
102 Val Gln Leu Gln Asp Asn Gly Tyr Asn Gly Leu Leu Ile Ala Ile Asn
103 35 40 45

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108 Phe Phe Arg Asn Ile Lys Ile Leu Ile Pro Ala Thr Trp Lys Ala Asn
109                      85                      90                      95
110 Asn Asn Ser Lys Ile Lys Gln Glu Ser Tyr Glu Lys Ala Asn Val Ile
111                      100                      105                      110
112 Val Thr Asp Trp Tyr Gly Ala His Gly Asp Asp Pro Tyr Thr Leu Gln
113                      115                      120                      125
114 Tyr Arg Gly Cys Gly Lys Glu Gly Lys Tyr Ile His Phe Thr Pro Asn
115                      130                      135                      140
116 Phe Leu Leu Asn Asp Asn Leu Thr Ala Gly Tyr Gly Ser Arg Gly Arg
117 145                      150                      155                      160
118 Val Phe Val His Glu Trp Ala His Leu Arg Trp Gly Val Phe Asp Glu
119                      165                      170                      175
120 Tyr Asn Asn Asp Lys Pro Phe Tyr Ile Asn Gly Gln Asn Gln Ile Lys
121                      180                      185                      190
122 Val Thr Arg Cys Ser Ser Asp Ile Thr Gly Ile Phe Val Cys Glu Lys
123                      195                      200                      205
124 Gly Pro Cys Pro Gln Glu Asn Cys Ile Ile Ser Lys Leu Phe Lys Glu
125                      210                      215                      220
126 Gly Cys Thr Phe Ile Tyr Asn Ser Thr Gln Asn Ala Thr Ala Ser Ile
127 225                      230                      235                      240
128 Met Phe Met Gln Ser Leu Ser Ser Val Val Glu Phe Cys Asn Ala Ser
129                      245                      250                      255
130 Thr His Asn Gln Glu Ala Pro Asn Leu Gln Asn Gln Met Cys Ser Leu
131                      260                      265                      270
132 Arg Ser Ala Trp Asp Val Ile Thr Asp Ser Ala Asp Phe His His Ser
133                      275                      280                      285
134 Phe Pro Met Asn Gly Thr Glu Leu Pro Pro Pro Pro Thr Phe Ser Leu
135                      290                      295                      300
136 Val Glu Ala Gly Asp Lys Val Val Cys Leu Val Leu Asp Val Ser Ser
137 305                      310                      315                      320
138 Lys Met Ala Glu Ala Asp Arg Leu Leu Gln Leu Gln Gln Ala Ala Glu
139                      325                      330                      335
140 Phe Tyr Leu Met Gln Ile Val Glu Ile His Thr Phe Val Gly Ile Ala
141                      340                      345                      350
142 Ser Phe Asp Ser Lys Gly Glu Ile Arg Ala Gln Leu His Gln Ile Asn
143                      355                      360                      365
144 Ser Asn Asp Asp Arg Lys Leu Leu Val Ser Tyr Leu Pro Thr Thr Val
145                      370                      375                      380
146 Ser Ala Lys Thr Asp Ile Ser Ile Cys Ser Gly Leu Lys Lys Gly Phe
147 385                      390                      395                      400
148 Glu Val Val Glu Lys Leu Asn Gly Lys Ala Tyr Gly Ser Val Met Ile
149                      405                      410                      415
150 Leu Val Thr Ser Gly Asp Asp Lys Leu Leu Gly Asn Cys Leu Pro Thr
151                      420                      425                      430
152 Val Leu Ser Ser Gly Ser Thr Ile His Ser Ile Ala Leu Gly Ser Ser

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156 Phe Phe Val Pro Asp Ile Ser Asn Ser Asn Ser Met Ile Asp Ala Phe
157 465          470          475          480
158 Ser Arg Ile Ser Ser Gly Thr Gly Asp Ile Phe Gln Gln His Ile Gln
159          485          490          495
160 Leu Glu Ser Thr Gly Glu Asn Val Lys Pro His His Gln Leu Lys Asn
161          500          505          510
162 Thr Val Thr Val Asp Asn Thr Val Gly Asn Asp Thr Met Phe Leu Val
163          515          520          525
164 Thr Trp Gln Ala Ser Gly Pro Pro Glu Ile Ile Leu Phe Asp Pro Asp
165          530          535          540
166 Gly Arg Lys Tyr Tyr Thr Asn Asn Phe Ile Thr Asn Leu Thr Phe Arg
167 545          550          555          560
168 Thr Ala Ser Leu Trp Ile Pro Gly Thr Ala Lys Pro Gly His Trp Thr
169          565          570          575
170 Tyr Thr Leu Asn Asn Thr His His Ser Leu Gln Ala Leu Lys Val Thr
171          580          585          590
172 Val Thr Ser Arg Ala Ser Asn Ser Ala Val Pro Pro Ala Thr Val Glu
173          595          600          605
174 Ala Phe Val Glu Arg Asp Ser Leu His Phe Pro His Pro Val Met Ile
175          610          615          620
176 Tyr Ala Asn Val Lys Gln Gly Phe Tyr Pro Ile Leu Asn Ala Thr Val
177 625          630          635          640
178 Thr Ala Thr Val Glu Pro Glu Thr Gly Asp Pro Val Thr Leu Arg Leu
179          645          650          655
180 Leu Asp Asp Gly Ala Gly Ala Asp Val Ile Lys Asn Asp Gly Ile Tyr
181          660          665          670
182 Ser Arg Tyr Phe Phe Ser Phe Ala Ala Asn Gly Arg Tyr Ser Leu Lys
183          675          680          685
184 Val His Val Asn His Ser Pro Ser Ile Ser Thr Pro Ala His Ser Ile
185          690          695          700
186 Pro Gly Ser His Ala Met Tyr Val Pro Gly Tyr Thr Ala Asn Gly Asn
187 705          710          715          720
188 Ile Gln Met Asn Ala Pro Arg Lys Ser Val Gly Arg Asn Glu Glu Glu
189          725          730          735
190 Arg Lys Trp Gly Phe Ser Arg Val Ser Ser Gly Gly Ser Phe Ser Val
191          740          745          750
192 Leu Gly Val Pro Ala Gly Pro His Pro Asp Val Phe Pro Pro Cys Lys
193          755          760          765
194 Ile Ile Asp Leu Glu Ala Val Lys Val Glu Glu Glu Leu Thr Leu Ser
195          770          775          780
196 Trp Thr Ala Pro Gly Glu Asp Phe Asp Gln Gly Gln Ala Thr Ser Tyr
197 785          790          795          800
198 Glu Ile Arg Met Ser Lys Ser Leu Gln Asn Ile Gln Asp Asp Phe Asn
199          805          810          815
200 Asn Ala Ile Leu Val Asn Thr Ser Lys Arg Asn Pro Gln Gln Ala Gly
201          820          825          830

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204 Glu His Gln Pro Asn Gly Glu Thr His Glu Ser His Arg Ile Tyr Val
205      850      855      860
206 Ala Ile Arg Ala Met Asp Arg Asn Ser Leu Gln Ser Ala Val Ser Asn
207 865      870      875      880
208 Ile Ala Gln Ala Pro Leu Phe Ile Pro Pro Asn Ser Asp Pro Val Pro
209      885      890      895
210 Ala Arg Asp Tyr Leu Ile Leu Lys Gly Val Leu Thr Ala Met Gly Leu
211      900      905      910
212 Ile Gly Ile Ile Cys Leu Ile Ile Val Val Thr His His Thr Leu Ser
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220 <213> ORGANISM: Homo sapiens
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226 gcgcagcagc agcagcagca gcagcagcag caggcgccgc agctgagacc ggcgggccgac 240
227 ggccagccct caggggggcg tcacaagtca gcgcccgaagc aagtcaagcg acagcgctcg 300
228 tcttcgcccg aactgatgcg ctgcaaagcg cggctcaact tcagcggctt tggctacagc 360
229 ctgccgcagc agcagccggc cgccgtggcg cgccgcaacg agcgcgagcg caaccgcgctc 420
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236 agcag 785
238 <210> SEQ ID NO: 4
239 <211> LENGTH: 236
240 <212> TYPE: PRT
241 <213> ORGANISM: Homo sapiens
243 <400> SEQUENCE: 4
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247      20      25      30
248 Ala Thr Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Gln
249      35      40      45
250 Ser Ala Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Ala Pro
251      50      55      60
252 Gln Leu Arg Pro Ala Ala Asp Gly Gln Pro Ser Gly Gly Gly His Lys
253 65      70      75      80
254 Ser Ala Pro Lys Gln Val Lys Arg Gln Arg Ser Ser Ser Pro Glu Leu

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date